## STIC Biotechnology Systems Branch

## RAW SEQUENCE LISTING ERROR REPORT

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**Application Serial Number:** 

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 01/10/06



**IFWP** 

RAW SEQUENCE LISTING DATE: 09/15/2006
PATENT APPLICATION: US/10/591,817 TIME: 09:07:34

Input Set : A:\Sequenc Listing 2006\_1487A.txt
Output Set: N:\CRF4\09152006\J591817.raw

3 <110> APPLICANT: Kyoto University
5 <120> TITLE OF INVENTION: Pharmaceutical composition comprising a CXCR3 inhibitor
7 <130> FILE REFERENCE: 665078

C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/591,817

C--> 9 <141> CURRENT FILING DATE: 2006-09-06

9 <150> PRIOR APPLICATION NUMBER: JP JP 2004-065612

10 <151> PRIOR FILING DATE: 2004-03-09

12 <160> NUMBER OF SEQ ID NOS: 34

14 <170 > SOFTWARE: PatentIn version 3.1

## Dres Not Comply Corrected Diskette Neededy **ERRORED SEQUENCES** 430 <210> SEQ ID NO: 431 <211> LENGTH 432 <212> TYPE: DNA 433 <213> ORGANISM: Artificial sequence 435 <220> FEATURE: 436 <223> OTHER INFORMATION: DNA primer for PCR 438 <400> SEQUENCE: 24 E--> 439 tgaattcaagc catgtacctt gaggtta, 478 <210> SEQ ID NO: 28 479 <211> LENGTH: 30 480 <212> TYPE: DNA 481 <213> ORGANISM: Artificial sequence are allowed are allowed are allowed are allowed and renter a server a serve 483 <220> FEATURE: 484 <223> OTHER INFORMATION: DNA probe for PCR 486 <400> SEQUENCE: 28 E--> 487 ctacccttgg acccagaggt tctttgagtc 499 <210> SEQ ID NO: 30 500 <211> LENGTH 2 501 <212> TYPE: (RNA) 502 <213 > ORGANISM: Artificial sequence 504 <220> FEATURE: 505 <223 > OTHER INFORMATION: siRNA 507 <400> SEQUENCE: 30 E--> 508 guggcaggug cccucuucat 511 <210> SEO ID NO: 31 512 <211> LENGTH: 21 513 <212> TYPE RNA 514 <213 > ORGANISM. Artificial sequence 516 <220> FEATURE: 517 <223> OTHER INFORMATION: siRNA

RAW SEQUENCE LISTING

DATE: 09/15/2006

PATENT APPLICATION: US/10/591,817

TIME: 09:07:34

Input Set : A:\Sequenc Listing 2006 1487A.txt Output Set: N:\CRF4\09152006\J591817.raw

Same error 519 <400> SEQUENCE: 31 E--> 520 ugaagagggc accugccact 532 <210> SEQ ID NO: 33 533 <211> LENGTH 21 534 <212> TYPE: RNA 535 <213> ORGANISM: Artificial sequence 537 <220> FEATURE: 538 <223> OTHER INFORMATION: siRNA 540 <400> SEQUENCE: 33 E--> 541 cgagagugac ucgugcugut t 544 <210> SEQ ID NO: 34 545 <211> LENGTH 2 546 <212> TYPE KNA 547 <213> ORGANISM: Artificial sequence 549 <220> FEATURE: 550 <223> OTHER INFORMATION: siRN 552 <400> SEQUENCE: 34 E--> 553 acagcacgag ucacucucgt t PIS use "W" instead OF "E". RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/15/2006 PATENT APPLICATION: US/10/591,817 TIME: 09:07:35

Input Set : A:\Sequenc Listing 2006\_1487A.txt
Output Set: N:\CRF4\09152006\J591817.raw

## Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:28; Line(s) 487

VERIFICATION SUMMARY

DATE: 09/15/2006

PATENT APPLICATION: US/10/591,817

TIME: 09:07:35

Input Set : A:\Sequenc Listing 2006\_1487A.txt
Output Set: N:\CRF4\09152006\J591817.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:439 M:254 E: No. of Bases conflict, LENGTH:Input:27 Counted:28 SEQ:24
L:439 M:252 E: No. of Seq. differs, <211> LENGTH:Input:27 Found:28 SEQ:24
L:487 M:254 E: No. of Bases conflict, LENGTH:Input:21 Counted:30 SEQ:28
L:508 M:321 E: (1) "t" not allowed in RNA Sequence, NUMBER OF INVALID 't' KEYS:2
L:520 M:321 E: (1) "t" not allowed in RNA Sequence, NUMBER OF INVALID 't' KEYS:2
L:541 M:321 E: (1) "t" not allowed in RNA Sequence, NUMBER OF INVALID 't' KEYS:2
L:553 M:321 E: (1) "t" not allowed in RNA Sequence, NUMBER OF INVALID 't' KEYS:2